SEQUENCE LISTING

<110> GOVERNMENT OF THE UNITED STATES OF AMERICA, REPRESENTED BY THE SECRETARY, DEPARTMENT OF HEALTH AND HUMAN SERVICES
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KOMIYA, TAKEFUMI

<120> COMPOSITIONS AND METHODS FOR INHIBITING TRANSLATION OF A CHIMERIC GENE

<130> 221749

<140> 10/698,070
<141> 2003-10-30

<160> 12

<170> PatentIn version 3.3

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<212> DNA
<213> Homo sapiens

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35 40 45

Arg Lys Gln Val Val Asn Leu Ser Pro Ala Asn Ser Lys Arg Pro Asn
50 55 60

Gly Phe Val Asp Asn Ser Phe Leu Asp Ile Lys Arg Ile Arg Val Gly
65 70 75 80

Glu Asn Leu Ser Ala Gly Gln Gly Gly Leu Gln Ile Asn Asn Gly Gln
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Ser Gln Ile Met Ser Gly Thr Leu Pro Met Ser Gln Ala Pro Leu Arg
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Lys Thr Asn Thr Leu Pro Ser His Thr His Ser Pro Gly Asn Gly Leu
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Leu Gln Glu Leu Phe Asn Glu Leu Thr Asn Ile Ser Val Pro Pro Met
            180                 185                 190

Ser Asp Leu Glu Leu Glu Asn Met Ile Asn Ala Thr Ile Lys Gln Asp
        195                 200                 205

Asp Pro Phe Asn Ile Asp Leu Gly Gln Gln Ser Gln Arg Ser Thr Pro
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Arg Pro Ser Leu Pro Met Glu Lys Ile Val Ile Lys Ser Glu Tyr Ser
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Pro Gly Leu Thr Gln Gly Pro Ser Gly Ser Pro Gln Leu Arg Pro Pro
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Ser Ala Gly Pro Ala Phe Ser Met Ala Asn Ser Ala Leu Ser Thr Ser
            260                 265                 270

Ser Pro Ile Pro Ser Val Pro Gln Ser Gln Ala Gln Pro Gln Thr Gly
        275                 280                 285

Ser Gly Ala Ser Arg Ala Leu Pro Ser Trp Gln Glu Val Ser His Ala
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Gln Gln Leu Lys Gln Ile Ala Ala Asn Arg Gln Gln His Ala Arg Met
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Gln Gln His Gln Gln Gln His Gln Pro Thr Asn Trp Ser Ala Leu Pro
                325                 330                 335

Ser Ser Ala Gly Pro Ser Pro Gly Pro Phe Gly Gln Glu Lys Ile Pro
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Ser Pro Ser Phe Gly Gln Gln Thr Phe Ser Pro Gln Ser Ser Pro Met
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Pro Gly Val Ala Gly Gly Ser Gly Gln Ser Lys Val Met Ala Asn Tyr
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Met Tyr Lys Ala Gly Pro Ser Ala Gln Gly Gly His Leu Asp Val Leu
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405 410 415

His Pro Ala Met Glu Pro Arg Gln Gly Asn Thr Lys Pro Leu Phe His
420 425 430

Phe Asn Ser Asp Gln Ala Asn Gln Gln Met Pro Ser Val Leu Pro Ser
435 440 445

Gln Asn Lys Pro Ser Leu Leu His Tyr Thr Gln Gln Gln Gln Gln Gln
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Gln Gln Gln Gln Gln Pro Ser Ser Gln Pro Ala Gln Ser Leu Pro Ser
530 535 540

Gln Pro Leu Leu Arg Ser Pro Leu Pro Leu Gln Gln Lys Leu Leu Leu
545 550 555 560

Gln Gln Met Gln Asn Gln Pro Ile Ala Gly Met Gly Tyr Gln Val Ser
565 570 575

Gln Gln Gln Arg Gln Asp Gln His Ser Val Val Gly Gln Asn Thr Gly
580 585 590

Pro Ser Pro Ser Pro Asn Pro Cys Ser Asn Pro Asn Thr Gly Ser Gly
595 600 605

Tyr Met Asn Ser Gln Gln Ser Leu Leu Asn Gln Gln Leu Met Gly Lys
610 615 620

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Leu Gln Gln Gln Met Leu Ala Asp Ala Glu Lys Ile Ala Pro Gln Asp
                645                 650                 655

Gln Ile Asn Arg His Leu Ser Arg Pro Pro Pro Asp Tyr Lys Asp Gln
            660                 665                 670

Arg Arg Asn Val Gly Asn Met Gln Pro Thr Ala Gln Tyr Ser Gly Gly
        675                 680                 685

Ser Ser Thr Ile Ser Leu Asn Ser Asn Gln Ala Leu Ala Asn Pro Val
    690                 695                 700

Ser Thr His Thr Ile Leu Thr Pro Asn Ser Ser Leu Leu Ser Thr Ser
705                 710                 715                 720

His Gly Thr Arg Met Pro Ser Leu Ser Thr Ala Val Gln Asn Met Gly
                725                 730                 735

Met Tyr Gly Asn Leu Pro Cys Asn Gln Pro Asn Thr Tyr Ser Val Thr
            740                 745                 750

Ser Gly Met Asn Gln Leu Thr Gln Gln Arg Asn Pro Lys Gln Leu Leu
        755                 760                 765

Ala Asn Gln Asn Asn Pro Met Met Pro Arg Pro Pro Thr Leu Gly Pro
    770                 775                 780

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785                 790                 795                 800

Ser Gln Gln Leu Arg Pro Asn Leu Thr His Ser Met Ala Ser Met Pro
                805                 810                 815

Pro Gln Arg Thr Ser Asn Val Met Ile Thr Ser Asn Thr Thr Ala Pro
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Asn Trp Ala Ser Gln Glu Gly Thr Ser Lys Gln Gln Glu Ala Leu Thr
        835                 840                 845

Ser Ala Gly Val Arg Phe Pro Thr Gly Thr Pro Ala Ala Tyr Thr Pro
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Arg Gly Leu Asn Leu Arg Pro Asn Gln Leu Ser Thr Gln Ile Leu Pro
        915                 920                 925

Asn Leu Asn Gln Ser Gly Thr Gly Leu Asn Gln Ser Arg Thr Gly Ile
    930                 935                 940

Asn Gln Pro Pro Ser Leu Thr Pro Ser Asn Phe Pro Ser Pro Asn Gln
945                 950                 955                 960

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Ser
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